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RAW SEQUENCE LISTING DATE: 02/03/2003 PATENT APPLICATION: US/09/920,262A TIME: 18:35:38 Input Set : A:\PTO.AMC.txt Output Set: N:\CRF4\02032003\I920262A.raw 4 <110> APPLICANT: Shealy, David Knight, David Scallon, Bernie Giles-Komar, Jill Peritt, David 10 <120> TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES, COMPOSITIONS, METHODS AND USES 12 <130> FILE REFERENCE: CEN0248 C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/920,262A C--> 15 <141> CURRENT FILING DATE: 2002-05-06 17 <150> PRIOR APPLICATION NUMBER: 60/223,358 18 <151> PRIOR FILING DATE: 2000-08-07 20 <150> PRIOR APPLICATION NUMBER: 60/236,827 21 <151> PRIOR FILING DATE: 2000-09-29 23 <160> NUMBER OF SEQ ID NOS: 15 25 <170> SOFTWARE: PatentIn Ver 3.1 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 5 29 <212> TYPE: PRT 30 <213> ORGANISM: Homo sapiens --> 31 <400> SEQUENCE: 1 33 Thr Tyr Trp Leu Gly 37 <210> SEQ ID NO: 2 38 <211> LENGTH: 17 39 <212> TYPE: PRT 40 <213> ORGANISM: Homo sapiens W--> 41 <400> SEQUENCE: 2 43 Ile Met Ser Pro Val Asp Ser Asp Ile Arg Tyr Ser Pro Ser Phe Gln 10 49 <210> SEQ ID NO: 3 50 <211> LENGTH: 10 51 <212> TYPE: PRT 52 <213> ORGANISM: Homo sapiens W--> 53 <400> SEQUENCE: 3 55 Pro Arg Pro Gly Gln Gly Tyr Phe Asp Phe

58 <210> SEQ ID NO: 4 59 <211> LENGTH: 11 60 <212> TYPE: PRT

W--> 62 <400> SEQUENCE: 4

61 <213> ORGANISM: Homo sapiens

64 Arg Ala Ser Gln Gly Ile Ser Ser Trp Leu Ala

6

7

34 1

44 1

46 Gly

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                      Input Set : A:\PTO.AMC.txt
                     Output Set: N:\CRF4\02032003\I920262A.raw
     65 1
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     67 <210> SEQ ID NO: 5
     68 <211> LENGTH: 7
     69 <212> TYPE: PRT
     70 <213> ORGANISM: Homo sapiens
W--> 71 <400> SEQUENCE: 5
     73 Ala Ala Ser Ser Leu Gln Ser
     76 <210> SEQ ID NO: 6
     77 <211> LENGTH: 9
     78 <212> TYPE: PRT
     79 <213> ORGANISM: Homo sapiens
W--> 80 <400> SEQUENCE: 6
     82 Gln Gln Tyr Asn Ile Tyr Pro Tyr Thr
     85 <210> SEQ ID NO: 7
     86 <211> LENGTH: 119
     87 <212> TYPE: PRT
     88 <213> ORGANISM: Homo sapiens
W--> 89 <400> SEQUENCE: 7
     91 Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
     94 Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Tyr
     97 Trp Leu Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Asp Trp Ile
                                    40
     100 Gly Ile Met Ser Pro Val Asp Ser Asp Ile Arg Tyr Ser Pro Ser Phe
                                 55
     103 Gln Gly Gln Val Thr Met Ser Val Asp Lys Ser Ile Thr Thr Ala Tyr
                             70
     106 Leu Gln Trp Asn Ser Leu Lys Ala Ser Asp Thr Ala Met Tyr Tyr Cys
                         85
     109 Ala Arg Arg Pro Gly Gln Gly Tyr Phe Asp Phe Trp Gly Gln Gly
     110
                     100
                                         105
     112 Thr Leu Val Thr Val Ser Ser
     113
                115
     115 <210> SEO ID NO: 8
     116 <211> LENGTH: 108
     117 <212> TYPE: PRT
     118 <213> ORGANISM: Homo sapiens
W--> 119 <400> SEQUENCE: 8
     121 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
     122 1
    124 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser Trp
    127 Leu Ala Trp Tyr Gln Gln Lys Pro Glu Lys Ala Pro Lys Ser Leu Ile
                35
                                     40
    130 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
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55

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133 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                         70
  136 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn Ile Tyr Pro Tyr
                    85
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  139 Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
        . 100
  142 <210> SEQ ID NO: 9
  143 <211> LENGTH: 503
  144 <212> TYPE: PRT
  145 <213> ORGANISM: Homo sapiens
-> 146 <400> SEQUENCE: 9
  148 Arg Asn Leu Pro Val Ala Thr Pro Asp Pro Gly Met Phe Pro Cys Leu
  151 His His Ser Gln Asn Leu Leu Arg Ala Val Ser Asn Met Leu Gln Lys
                20
                                     25
  154 Ala Arg Gln Thr Leu Glu Phe Tyr Pro Cys Thr Ser Glu Glu Ile Asp
            35
  157 His Glu Asp Ile Thr Lys Asp Lys Thr Ser Thr Val Glu Ala Cys Leu
                             5.5
  160 Pro Leu Glu Leu Thr Lys Asn Glu Ser Cys Leu Asn Ser Arg Glu Thr
                        70
  163 Ser Phe Ile Thr Asn Gly Ser Cys Leu Ala Ser Arg Lys Thr Ser Phe
 166 Met Met Ala Leu Cys Leu Ser Ser Ile Tyr Glu Asp Leu Lys Met Tyr
                 100
                                     105
 169 Gln Val Glu Phe Lys Thr Met Asn Ala Lys Leu Leu Met Asp Pro Lys
             115
                                 120
 172 Arg Gln Ile Phe Leu Asp Gln Asn Met Leu Ala Val Ile Asp Glu Leu
 173 130
                             135
                                                140
 175 Met Gln Ala Leu Asn Phe Asn Ser Glu Thr Val Pro Gln Lys Ser Ser
                         150
                                             155
 178 Leu Glu Glu Pro Asp Phe Tyr Lys Thr Lys Ile Lys Leu Cys Ile Leu
 179 . 165
                                        170
 181 Leu His Ala Phe Arg Ile Arg Ala Val Thr Ile Asp Arg Val Met Ser
                180
                                    185
 184 Tyr Leu Asn Ala Ser Ile Trp Glu Leu Lys Lys Asp Val Tyr Val Val
            195
                                200
 187 Glu Leu Asp Trp Tyr Pro Asp Ala Pro Gly Glu Met Val Val Leu Thr
                             215
 190 Cys Asp Thr Pro Glu Glu Asp Gly Ile Thr Trp Thr Leu Asp Gln Ser
                         230
                                            235
 193 Ser Glu Val Leu Gly Ser Gly Lys Thr Leu Thr Ile Gln Val Lys Glu
                    245
                                        250
 196 Phe Gly Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly Glu Val Leu
                                    265
 199 Ser His Ser Leu Leu Leu Leu His Lys Lys Glu Asp Gly Ile Trp Ser
      275
                                280
 202 Thr Asp Ile Leu Lys Asp Gln Lys Glu Pro Lys Asn Lys Thr Phe Leu
 203
                            295
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205 Arg Cys Glu Ala Lys Asn Tyr Ser Gly Arg Phe Thr Cys Trp Trp Leu
                             310
                                                  315
     208 Thr Thr Ile Ser Thr Asp Leu Thr Phe Ser Val Lys Ser Ser Arg Gly
                         325
                                              330
     211 Ser Ser Asp Pro Gln Gly Val Thr Cys Gly Ala Ala Thr Leu Ser Ala
                     340
                                          345
     214 Glu Arg Val Arg Gly Asp Asn Lys Glu Tyr Glu Tyr Ser Val Glu Cys
                                      360
     217 Gln Glu Asp Ser Ala Cys Pro Ala Ala Glu Glu Ser Leu Pro Ile Glu
                                 375
     220 Val Met Val Asp Ala Val His Lys Leu Lys Tyr Glu Asn Tyr Thr Ser
     221 385
                             390
                                                 395
     223 Ser Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro Lys Asn Leu
                         405
                                             410
     226 Gln Leu Lys Pro Leu Lys Asn Ser Arg Gln Val Glu Val Ser Trp Glu
                     420
                                         425
     229 Tyr Pro Asp Thr Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Thr Phe
     230 435
                                     440
     232 Cys Val Gln Val Gln Gly Lys Ser Lys Arg Glu Lys Lys Asp Arg Val
     233 450
                                 455
                                                     460
     235 Phe Thr Asp Lys Thr Ser Ala Thr Val Ile Cys Arg Lys Asn Ala Ser
                             470
                                                 475
     238 Ile Ser Val Arg Ala Gln Asp Arg Tyr Tyr Ser Ser Ser Trp Ser Glu
                         485
     241 Trp Ala Ser Val Pro Cys Ser
     242
                     500
     244 <210> SEQ ID NO: 10
     245 <211> LENGTH: 15
     246 <212> TYPE: DNA
     247 <213> ORGANISM: Homo sapiens
W--> 248 <400> SEQUENCE: 10
     250 agatatacta tgcac
                                                                           15
     252 <210> SEQ ID NO: 11
     253 <211> LENGTH: 51
     254 <212> TYPE: DNA
     255 <213> ORGANISM: Homo sapiens
W--> 256 <400> SEQUENCE: 11
     258 gttatatcat ttgatggaag caataaatac tacgtagact ccgtgaaggg c
                                                                           51
     260 <210> SEQ ID NO: 12
     261 <211> LENGTH: 30
     262 <212> TYPE: DNA
     263 <213> ORGANISM: Homo sapiens
W--> 264 <400> SEQUENCE: 12
     266 gaggcccggg gatcgtatgc ttttgatatc
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     269 <211> LENGTH: 42
     270 <212> TYPE: DNA
     271 <213> ORGANISM: Homo sapiens
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	274	ctctcctqca qqqccaqtca qaqtqttaqc aqctacttaq cc	42
		<210> SEQ ID NO: 14	
	277	<211> LENGTH: 18	
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	282	gatgcatcca acagggcc	18
	284	<210> SEQ ID NO: 15	
	285	<211> LENGTH: 21	
	286	<212> TYPE: DNA	
	287	<213> ORGANISM: Homo sapiens	
W>	288	<400> SEQUENCE: 15	
	290	cagcagcgta gcaactggcc t	21

VERIFICATION SUMMARY

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L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:31 M:283 W: Missing Blank Line separator, <400> field identifier
L:41 M:283 W: Missing Blank Line separator, <400> field identifier
L:53 M:283 W: Missing Blank Line separator, <400> field identifier
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L:80 M:283 W: Missing Blank Line separator, <400> field identifier
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L:146 M:283 W: Missing Blank Line separator, <400> field identifier
L:248 M:283 W: Missing Blank Line separator, <400> field identifier
L:256 M:283 W: Missing Blank Line separator, <400> field identifier
L:264 M:283 W: Missing Blank Line separator, <400> field identifier
L:272 M:283 W: Missing Blank Line separator, <400> field identifier
L:280 M:283 W: Missing Blank Line separator, <400> field identifier
L:288 M:283 W: Missing Blank Line separator, <400> field identifier
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